# Data Profiling and Data Quality Checks in SAS

Using SAS Studio on SAS On Demand for Academics (SODA)



### Imported Your Data Already?

 If you already have your data in SAS Studio on SAS On Demand for Academics, you can skip the slides providing an overview of the import process.



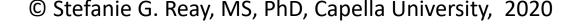


#### Dataset

• This tutorial is a walkthrough with a sample set of data. You may use this to walk through the tutorial, if you wish, but for your assignments, you will be asked to use your own dataset (as specified within the course).

#### Dataset reference:

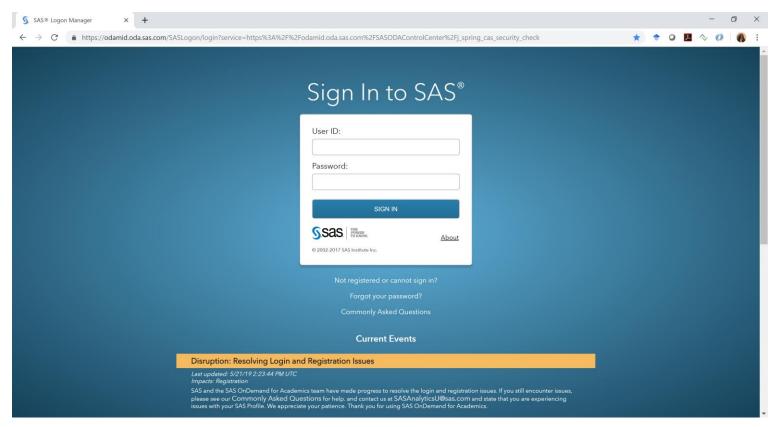
Skoryk, M. (2021). Sepsis Prediction from Clinical Data. Version 1. Retrieved from https://www.kaggle.com/maxskoryk/datasepsis





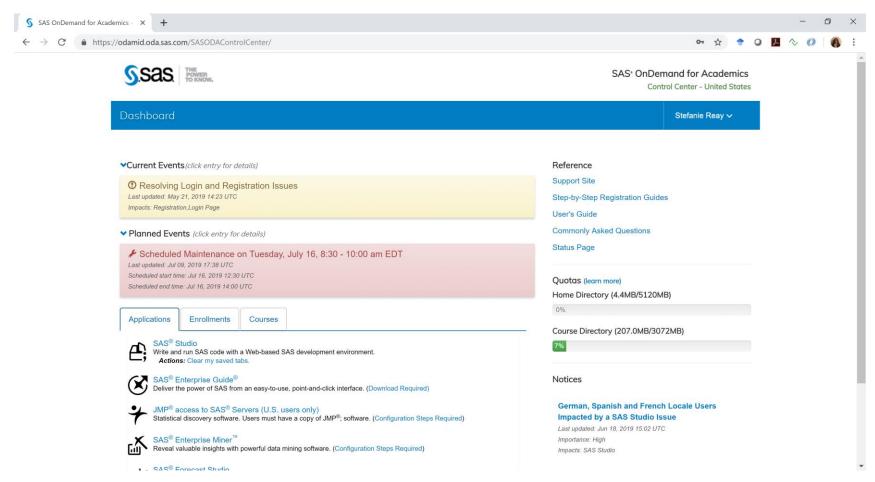
### Access the SAS OnDemand for Academics Control Center

#### https://odamid.oda.sas.com/SASODAControlCenter





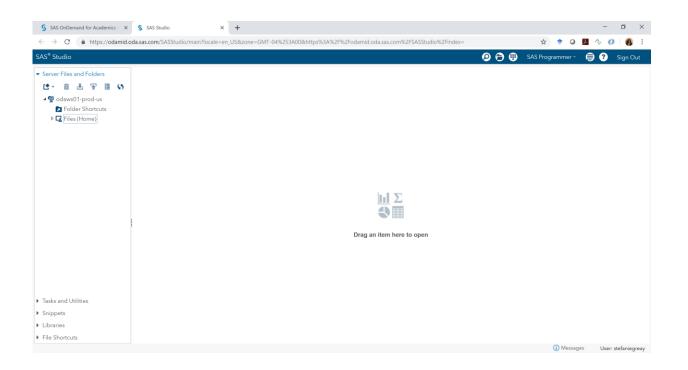
### SAS OnDemand for Academics (SODA) Control Center





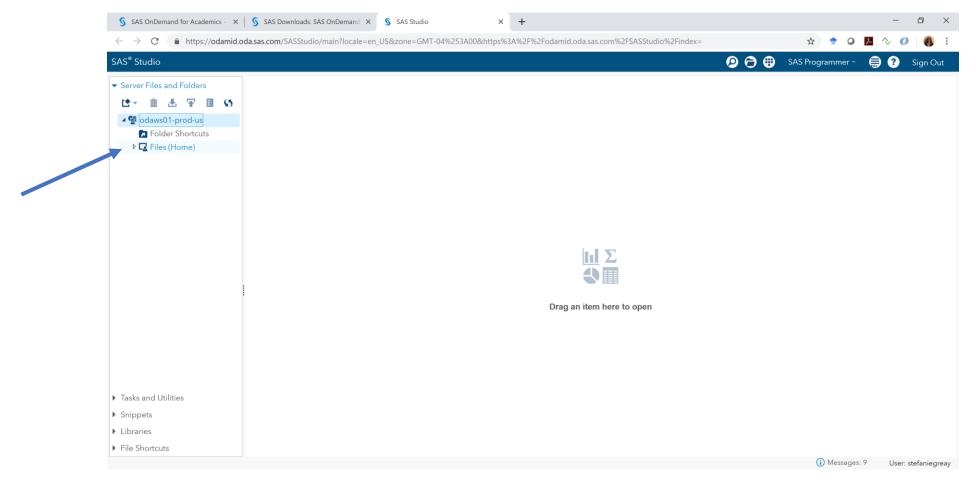
#### SAS Studio

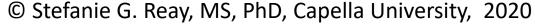
#### https://odamid.oda.sas.com/SASStudio/





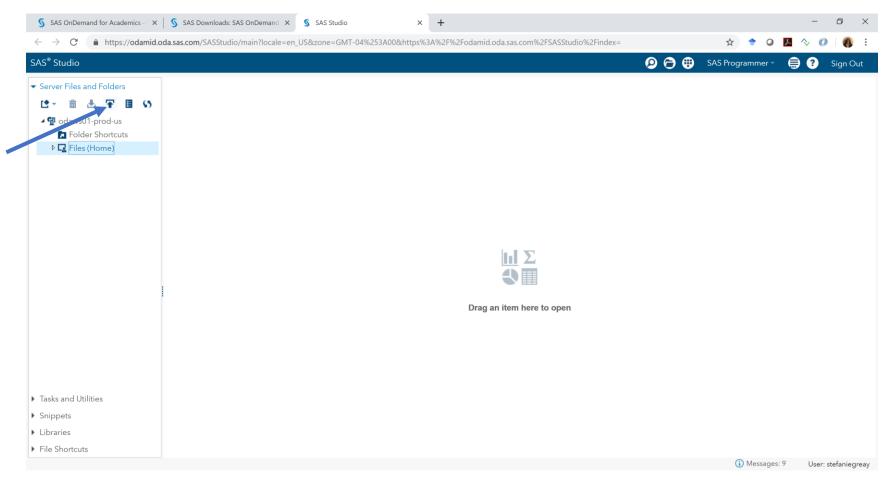
### Click on Files(Home)





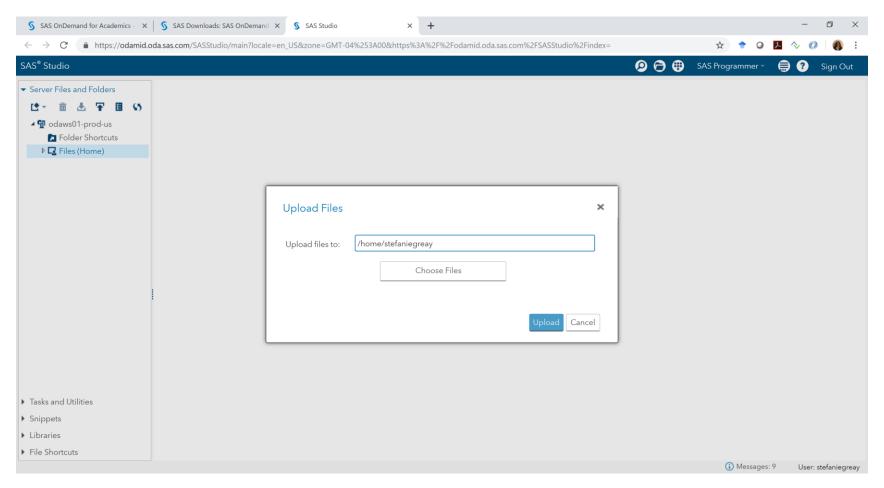


### The Upload button will display in dark blue



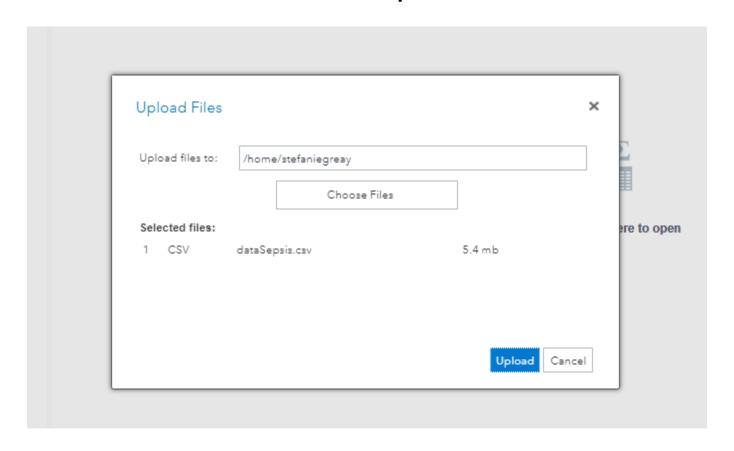


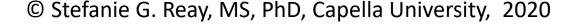
### You can create a folder at this point, if you wish, or simply upload to your home directory.





Select "Choose Files" to browse your computer for the dataset you want to upload. Once the dataset has been selected, click "Upload."

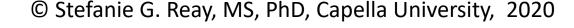






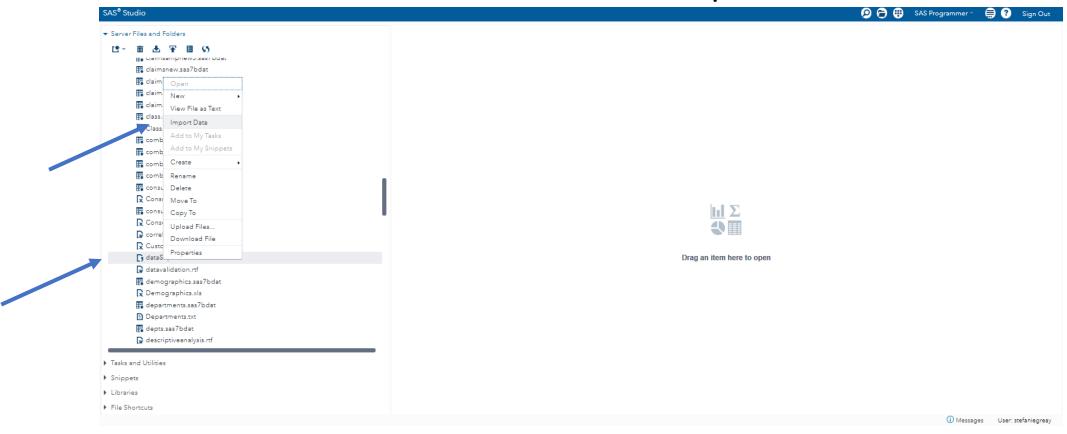
You will be able to view your files by clicking on "Files(Home)" to verify that your file successfully uploaded.

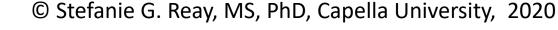






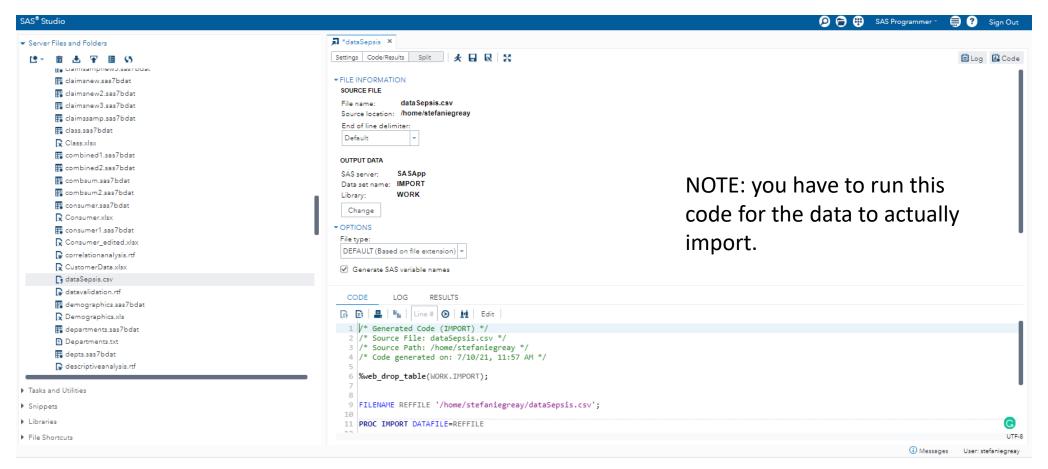
To import the dataset into a SAS dataset format (from the current csv format), right click on the name of the file, and select "Import Data."





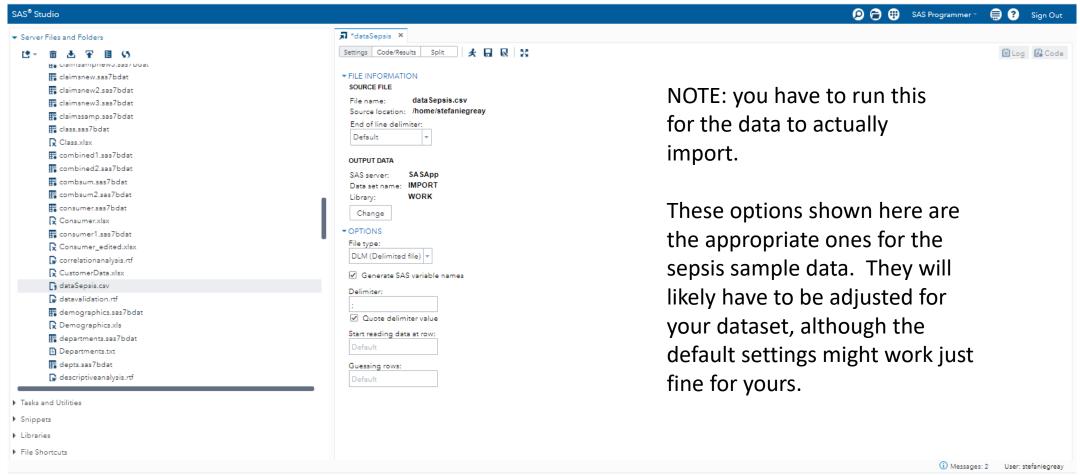


### The Proc Import code will be written for you (save this as a template to use for future imports!)



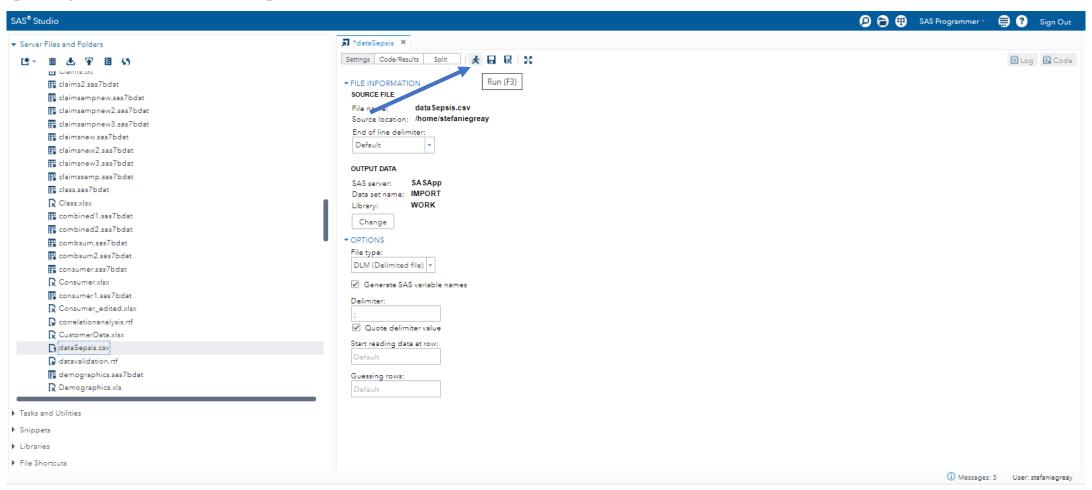


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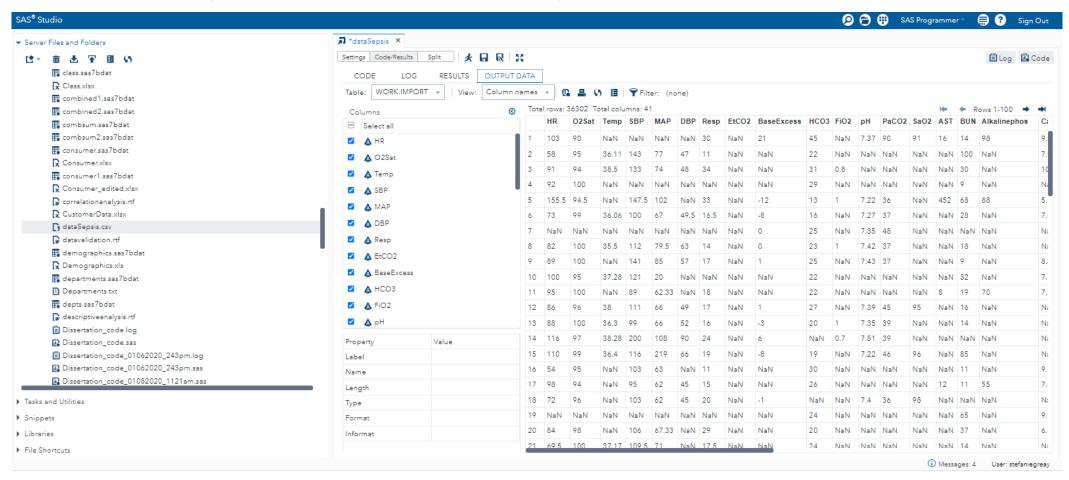
# To run the code, click the icon that looks like a guy running.





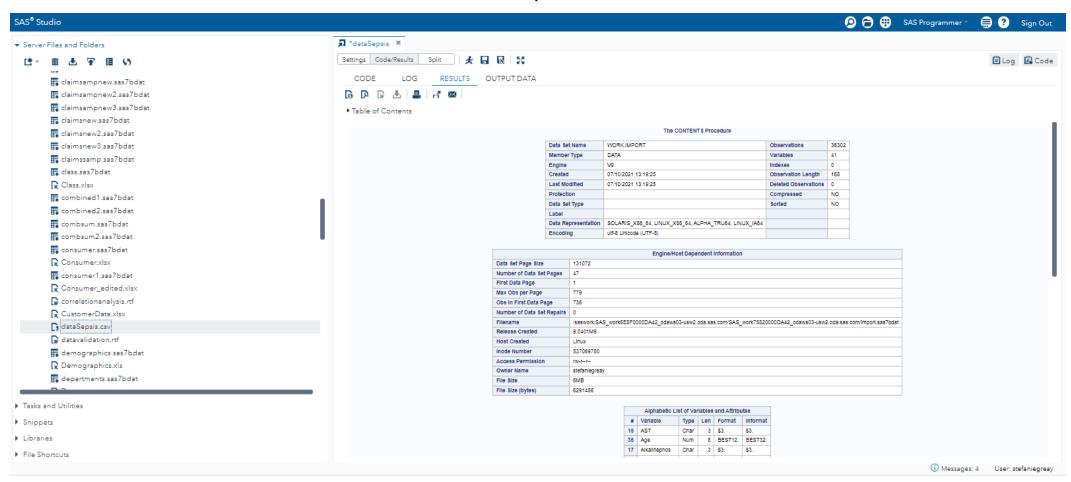


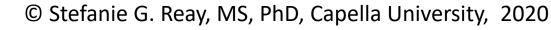
When you run the import, you will see the dataset and summary in the output data window when you click the "Code/Results" or "Split" tab and then "Output Data" and can verify its success.





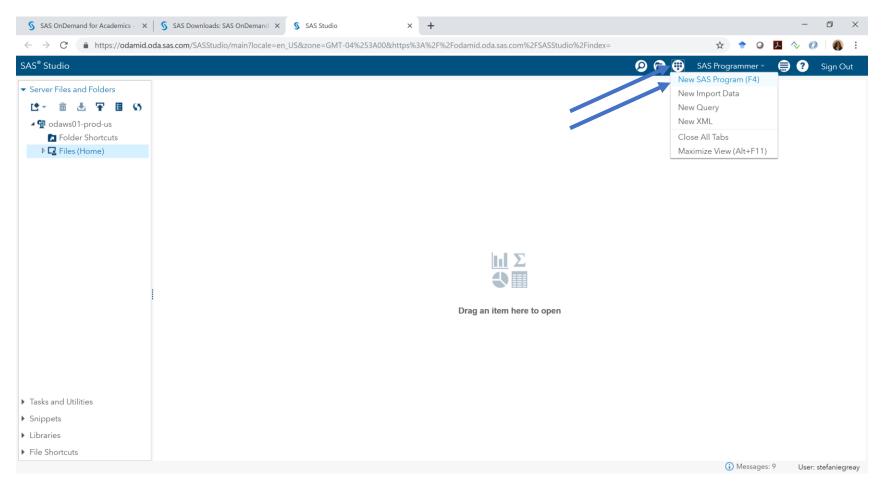
When you click the "Code/Results" or "Split" tab and then "Results," you can see the contents of the dataset, to verify the number of observations and variables are as expected.





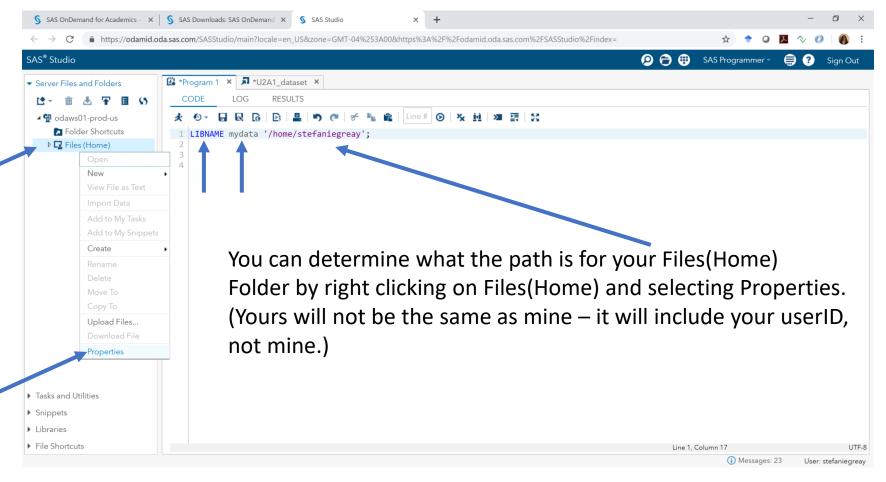


# To get started working with the dataset you just imported, start a new SAS program.



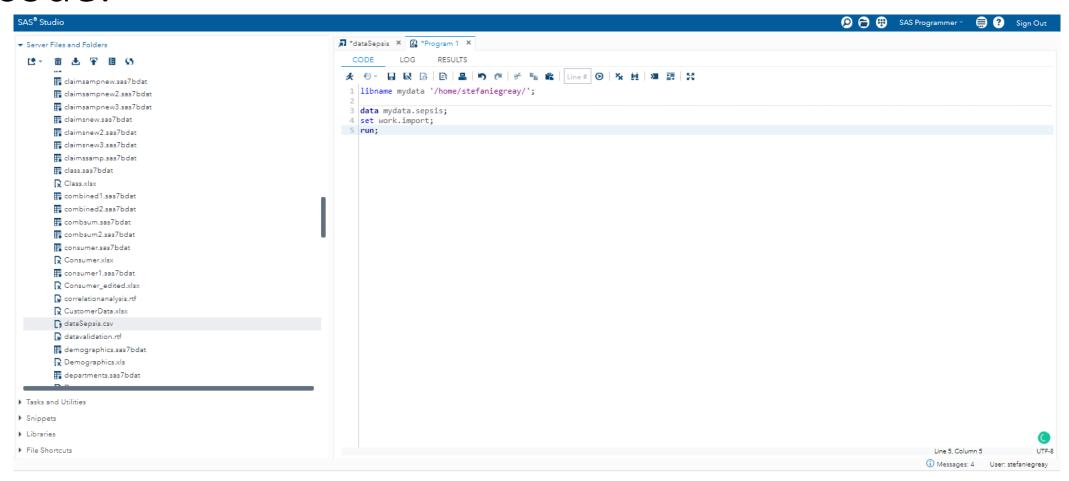


# To create a SAS Library for your Files (Home) folder, you need to use a libname statement





Save the temporary SAS dataset created by the import to your library using the following sample code.







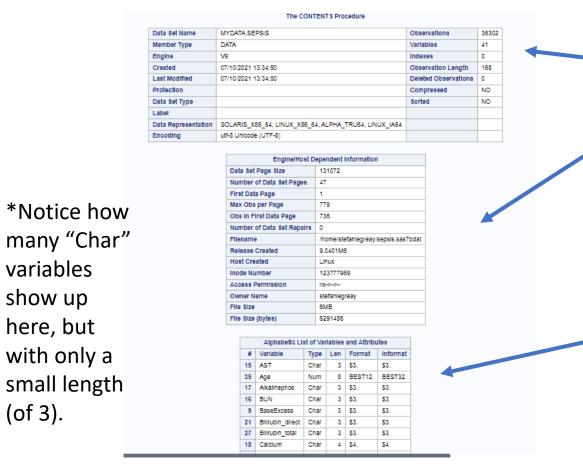
#### First: Look at the contents of the dataset

Proc contents provides the contents of the dataset identified next to the data= option. Remember that the part that comes before the . Is the library reference (SAS's reference for the folder) and the part that comes after is the name of the actual dataset.

Proc print prints the dataset out to the results/output window. We don't want to do this without limiting the number of rows output, though, because it could crash if we have a very large dataset. We can limit the number of records/rows printed using the obs= option in parentheses. This example prints the first 25 rows.



#### Proc Contents Output



Basic dataset information is here, including the number of observations (rows) and variables (columns).

The details of all of the variables and attributes of the variables are shown here, including the variable name, variable number (order of the variable in the input file), type (only Character (Char) or Numeric (Num) variable types exist in SAS), length, format (how SAS displays the data), and informat (how the data was read in from the input file).

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(of 3).



### Proc Print Output

Variables are across the top (each column is a variable).

	Obs	HR	O2 Sat	Temp	SBP	MAP	DBP	Resp	EtCO2	BaseExcess	HCOS	FIO2	рН	PaCO2	\$802	AST	BUN	Alkalinephos	Calcium	Chloride	Creatinine	Bilirubin direct	Glucose	Lactate	Magnesium	Phosphate	Potassium	BII
	1	103	90	NaN	NaN	NaN	NaN	30	NaN	21	45	NaN	7.37	90	91	16	14	02	0.3	95	0.7	MaM	103	MaN	2	3.3	3.8	0.3
	2	58	95	36.11	143	77	47	11	NaN	NaN	22	NaN	NaN	NaN	NaN	NaN	100	Each c	all ro	nro	contc	tha val		f+hc			5.1	Na
	3	91	94	38.5	133	74	48	34	NaN	NaN	31	0.8	NaN	NaN	NaN	NaN	30	Each	enre	pres	ents	the val	ue o	ı tile	;		3.8	N
	4	92	100	NaN	NaN	Naiv	MeN	NaN	NaN	NaN	29	NaN	NaN	NaN	NaN	NaN	9	variable identified in the column header									3.8	1
	5	155.5	94.5	NaN	147.5	102	NaN	33	NaN		13	1	7.22	36	NaN	452	68	- variable lacifulied in the column header									4.6	
	6	73	99	36.06	100	67	49.5	16.5	NaN	-8	16	Weis	7.97	37	NaN	NaN	28 -	for the	obs	erva	tion i	dentifie	ed or	า the	row		4.5	
	7	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	0	25	NaN	7.35	48	Nan	Ti-N	Na								_		4	1
	8	82	100	35.5	112	79.5	63	14	NaN	0	23	1	7.42		NaN	NaN	18	header/label. This cell, for example,								3.9	1	
	9	89	100	NaN	141	85	57	17	NaN	1	25	NaN	7.43		NaN	NaN	9	represents an O2Sat of 94 for observation								3.5	1	
	10		95	37.28	121	20	NaN	NaN	NaN	NaN	22	NaN	NaN		NaN	NaN	32									3.9	+	
		95	100	NaN	89	62.33	NaN	18	NaN	NaN	22	NaN	NaN	NaN	NaN	8	19										4.1	4
	12	-	96	38	111	66	49	17	NaN	1	27	NaN	7.39	45	95	NaN NaN	16	5 (the till d observation).										+
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	15		99	36.4	116	219	66	19	NaN	-8	19	NaN	7.22	46	96		85	NaN	NaN	96	8.7	NaN	74	NaN	1.9	NaN	4.9	+
	16		95	NaN	103	63	NaN	11	NaN	NaN	30	NaN	NaN	NaN	NaN	NaN		NaN	9.3	102	1	NaN	108	NaN	2.3	4.3	4.5	+
	17	_	94	NaN	95	62	45	15	NaN	NaN	26	NaN	NaN	NaN	NaN	12	11	55	7.4	101	0.5	NaN	122	NaN	2	2.9	4.1	t
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obs	erv	vat	ion)	•								NaN	NaN	NaN	NaN	NaN	14	NaN	NaN	102	0.9	NaN	96	NaN	2.4	NaN	3.7	Ť
	22	03	30	INDIN	10.0	00	40	14	INDIN	INDIN	20	NaN	NaN	NaN	NaN	NaN	19	NaN	8.3	109	0.9	NaN	140	NaN	1.7	4.2	4.3	
	23	88	96	NaN	97	65	45	28	NaN	-3	NaN	NaN	7.36	37	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	1.2	NaN	NaN	NaN	I
	24	80	99	NaN	129	100	77	18	NaN	NaN	22	NaN	NaN	NaN	NaN	NaN	8	NaN	8	107	0.6	NaN	118	NaN	1.8	2.8	3.8	+
	25	79	99	37.39	133	76	50	15	NaN	NaN	21	NaN	NaN	NaN	NaN	15	12	126	8.4	100	0.7	NaN	142	NaN	2	3.5	4.3	1

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## Second: Summarize the numeric variables and check for missing values Proc univariate outputs seven

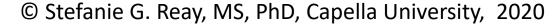
```
12
13 proc univariate data=mydata.sepsis plots;
14 run;
15
16 proc means data=mydata.sepsis nmiss;
17 run;
18
```

Sample code with var statement (to specify one or more specific numeric variables:

```
Proc univariate data=mydata.sepsis plots;
Var age;
Run;
```

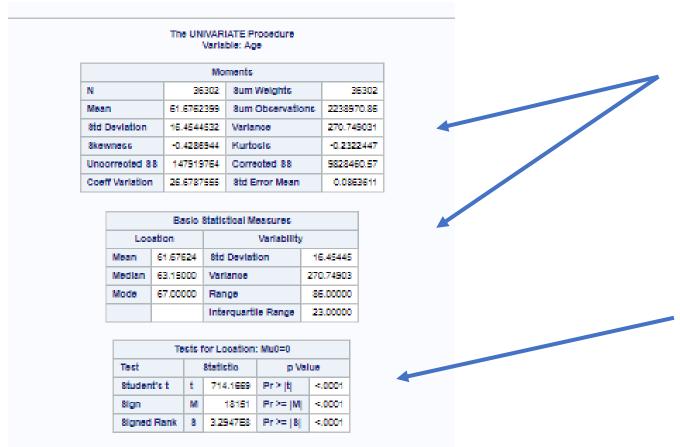
Proc univariate outputs several numeric summaries of the numeric variables, and the plots option adds several plots (the distribution plot, box plot, and normal quantile plot) for each variable. Not using a var statement tells SAS to do this summary for all variables with the type of Num (i.e. numeric variables).

Proc means with the nmiss option outputs the number of missing values for the numeric variables. Not using a var statement tells SAS to do this summary for all variables with the type of Num (i.e. numeric variables).





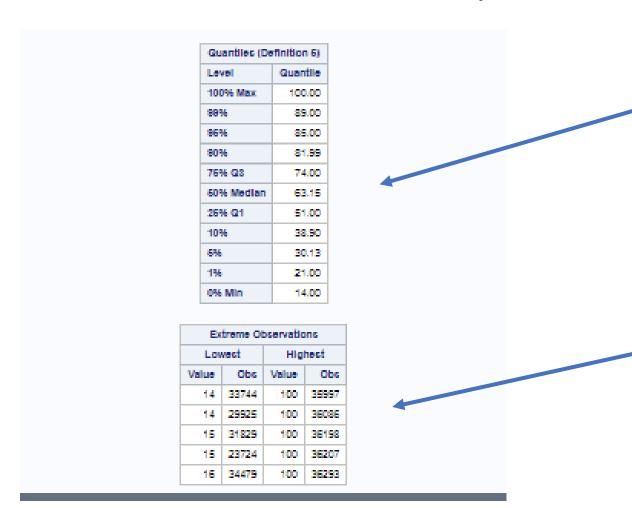
#### Proc Univariate Output



Basic summary statistics (measures of central tendency/middle; measures of dispersion/spread; distribution measures (skewness; kurtosis; etc.))

This section includes hypothesis testing details for testing if the middle of the distribution is centered around 0. For most data quality checks and data profiling, this section is irrelevant.

### Proc Univariate Output (Cont.)



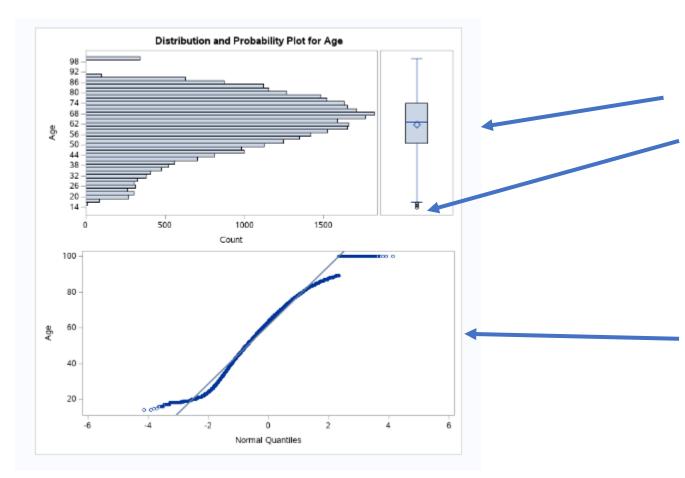
Percentile cutoff information for the variable is included here. For example, the 75<sup>th</sup> percentile is 74.00 in this case; the maximum is 100, and the minimum is 14.

Extreme observations are identified in this section. It includes the variable values themselves (for the lowest and highest observations), as well as the number identifying the observation (i.e. row) that that value can be found in. This is a great resource for identifying data that might have been entered incorrectly, like if someone entered 1000 for their age, instead of 100, for example.



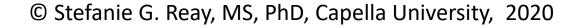


#### Proc Univariate Output (Cont.) - Plots



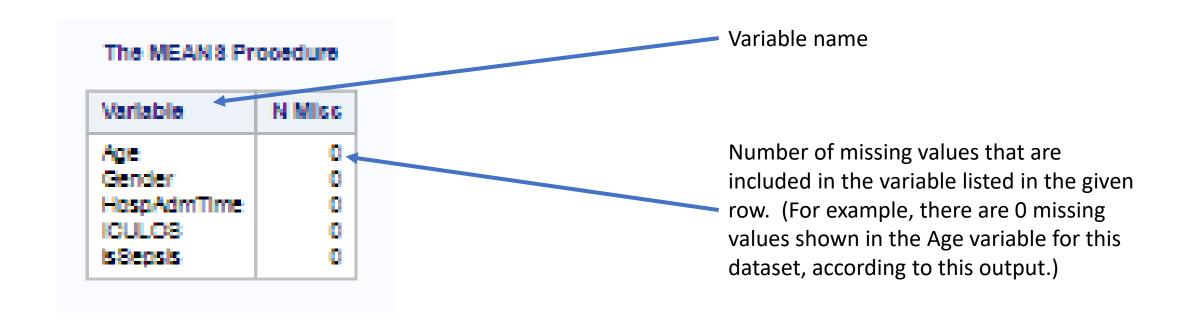
This section includes the histogram and box plot. Checking the overall shape of the distribution on these plots is a good step, as well as identifying any values that look out of place or like outliers (like the dots at the bottom of the box plot, for example).

The normal quantiles plot gives us an idea of whether or not the distribution is bell-shaped (or distributed according to a Normal distribution). The closer to all of the points following the lines, the closer to a perfect bell shape the distribution follows. Any points that stand far out from the rest may indicate ones that need to be explored further.





### Proc Means (with nmiss option) Output





<sup>\*</sup>Notice that there are only 5 variables that SAS identified as numeric and none of them have missing values, out of the 41 total variables in this dataset.

# Third: Summarize the character variables and check for missing values Proc freq outputs a table that counts the frequency (number of observations) with

```
18 | proc freq data=mydata.sepsis; 20 run; 21
```

Sample code with tables statement (to specify only using the variables formatted/recognized as character):

```
Proc freq data=mydata.sepsis;
Tables _CHAR_;
Run;
```

(To specify a specific variable, replace \_CHAR\_ with the name of the specific variable.)

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Proc freq outputs a table that counts the frequency (number of observations) with a particular value of that variable, as well as the relative frequency (or percent of observations), cumulative frequency and cumulative relative frequency). Not using a tables statement tells SAS to do this summary for all variables. This will cause a very large set of output if we don't specify at least character values only (example to the left).

You can technically also specify a numeric variable in the tables statement, although including continuous, numeric variables (ones with a lot of different values) is not beneficial or interpretable and will cause memory issues or crash the program if the data is very large. A variable like gender, here, however, that only has values of 0 and 1 (but was imported by SAS as numeric because of the 0's and 1's would be fine to include in a proc freq.



### Proc Freq Output Example

Variable value

Gender Frequency Percent Frequency Percent

1 20306 55.94 36302 100.00

Code used to get this output:

Proc freq data=mydata.sepsis; Tables gender; Run; Cumulative count and percent of observations with that variable value

Count of observations with that variable value (i.e. a Gender of 0 in this case)

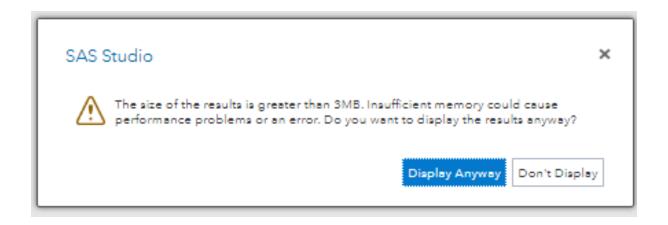
Percent of observations with that variable value (i.e. a Gender of 0 in this case)

A frequency table can be helpful for identifying variable values that look out of place (i.e have very small frequencies compared to the others), any spelling errors or differences in case (Fred and fred would be treated differently within SAS, as two different variable values, for example, because of the difference in capitalization, without further intervention).

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### Proc Freq for Variables with Many Values



This message results from the proc freq on this sepsis dataset without a tables statement. (It is trying to output one table per variable that SAS identified as character, with one row per unique variable value in each table.)

\*Note that this is not a common message and indicates that something might not be quite right with the data or how it was imported or how the variable types were recognized and registered by SAS.



### Code template

```
libname mydata '/home/stefaniegreay/';
                                                               proc univariate data=mydata.sepsis plots;
                                                               run;
data mydata.sepsis;
set work.import;
                                                               proc means data=mydata.sepsis nmiss;
run;
                                                               run;
                                                               proc freq data=mydata.sepsis;
proc contents data=mydata.sepsis;
                                                               tables gender;
run;
                                                               run;
proc print data=mydata.sepsis(obs=25);
                                                               proc freq data=mydata.sepsis;
run;
                                                               tables CHAR;
                                                               run;
```

